

OM nucleic - nucleic search, using sw model

Run on: October 6, 2005, 08:37:07 ; Search time 24747 Seconds  
(without alignments)  
11767.723 Million cell updates/sec

Title: US-10-616-624-1  
Perfect score: 6010  
Sequence: 1 gaattccagataggtagtc.....caggaagccccagcgccgg 6010

Scoring table: IDENTITY\_NUC  
Gapop 10.0 , Gapext 1.0

Searched: 4708233 seqs, 24227607955 residues

Total number of hits satisfying chosen parameters: 9416466

Minimum DB seq length: 0  
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%  
Maximum Match 100%  
Listing first 45 summaries

Database : GenEmbl:\*  
1: gb\_ba:\*  
2: gb\_htg:\*  
3: gb\_in:\*  
4: gb\_om:\*  
5: gb\_ov:\*  
6: gb\_pat:\*  
7: gb\_ph:\*  
8: gb\_pl:\*  
9: gb\_pr:\*  
10: gb\_ro:\*  
11: gb\_sts:\*  
12: gb\_sy:\*  
13: gb\_un:\*  
14: gb\_vi:\*

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Run on: October 6, 2005, 06:24:03 ; Search time 2855 Seconds  
(without alignments)  
12461.522 Million cell updates/sec

Title: US-10-616-624-1  
Perfect score: 6010  
Sequence: 1 gaattccagataggtagtc.....caggaagccccagcgccgg 6010

Scoring table: IDENTITY\_NUC  
Gapop 10.0 , Gapext 1.0

Searched: 4390206 seqs, 2959870667 residues

Total number of hits satisfying chosen parameters: 8780412

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%  
Maximum Match 100%  
Listing first 45 summaries

Database : N\_Geneseq\_16Dec04:\*  
1: geneseqn1980s:\*  
2: geneseqn1990s:\*  
3: geneseqn2000s:\*  
4: geneseqn2001as:\*  
5: geneseqn2001bs:\*  
6: geneseqn2002as:\*  
7: geneseqn2002bs:\*  
8: geneseqn2003as:\*  
9: geneseqn2003bs:\*  
10: geneseqn2003cs:\*  
11: geneseqn2003ds:\*  
12: geneseqn2004as:\*  
13: geneseqn2004bs:\*

GenCore version 5.1.6  
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Run on: October 6, 2005, 10:19:42 ; Search time 17440 Seconds  
(without alignments)  
13117.331 Million cell updates/sec

Title: US-10-616-624-1  
Perfect score: 6010  
Sequence: 1 gaattccagataggtagtcc.....caggaagccccagcgtccgg 6010

Scoring table: IDENTITY\_NUC  
Gapop 10.0 , Gapext 1.0 .

Searched: 34239544 seqs, 19032134700 residues

Total number of hits satisfying chosen parameters: 68479088

Minimum DB seq length: 0  
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%  
Maximum Match 100%  
Listing first 45 summaries

Database : EST:\*  
1: gb\_est1:\*  
2: gb\_est2:\*  
3: gb\_htc:\*  
4: gb\_est3:\*  
5: gb\_est4:\*  
6: gb\_est5:\*  
7: gb\_est6:\*  
8: gb\_gss1:\*  
9: gb\_gss2:\*

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OM protein - protein search, using sw model